SEQUENCE LISTING

<110> Johanson, Urban
West, Joanne
Dean, Caroline

<120> Arabidopsis thaliana derived Frigida gene conferring late flowering

<130> Mewburn

<140> US 09/890,475
<141> 2001-08-01

<150> PCT/GB00/00197
<151> 2000-01-25

<150> GB 9902660.1
<151> 1999-02-05

<160> 58

<170> PatentIn Ver. 2.1

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<223> Description of Artificial Sequence: FRI amino acid sequence predicted from the cDNA sequence

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Pro Lys Ile Val Glu Thr Glu Ser Thr Ser Met Asp Ile Thr Ile Gly
        35                  40                  45

Gln Ser Lys Gln Pro Gln Phe Leu Lys Ser Ile Asp Glu Leu Ala Ala
    50                  55                  60

Phe Ser Val Ala Val Glu Thr Phe Lys Arg Gln Phe Asp Asp Leu Gln
65                  70                  75                  80

Lys His Ile Glu Ser Ile Glu Asn Ala Ile Asp Ser Lys Leu Glu Ser
                85                  90                  95

Asn Gly Val Val Leu Ala Ala Arg Asn Asn Asn Phe His Gln Pro Met
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Leu Ser Pro Pro Arg Asn Asn Val Ser Val Glu Thr Thr Val Thr Val
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Ile Tyr Ala Asn Ile Ser Asp Gln Ala Lys Leu Met Glu Glu Ile Pro
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Ser Ala Leu Lys Leu Ala Lys Glu Pro Ala Lys Phe Val Leu Asp Cys
            180                 185                 190

Ile Gly Lys Phe Tyr Leu Gln Gly Arg Arg Ala Phe Thr Lys Glu Ser
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Pro Met Ser Ser Ala Arg Gln Val Ser Leu Leu Ile Leu Glu Ser Phe
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Leu Leu Met Pro Asp Arg Gly Lys Gly Lys Val Lys Ile Glu Ser Trp
225                 230                 235                 240

Ile Lys Asp Glu Ala Glu Thr Ala Ala Val Ala Trp Arg Lys Arg Leu
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Met Thr Glu Gly Gly Leu Ala Ala Ala Glu Lys Met Asp Ala Arg Gly
            260                 265                 270

Leu Leu Leu Leu Val Ala Cys Phe Gly Val Pro Ser Asn Phe Arg Ser
        275                 280                 285

Thr Asp Leu Leu Asp Leu Ile Arg Met Ser Gly Ser Asn Glu Ile Ala
    290                 295                 300

Gly Ala Leu Lys Arg Ser Gln Phe Leu Val Pro Met Val Ser Gly Ile
305                 310                 315                 320

Val Glu Ser Ser Ile Lys Arg Gly Met His Ile Glu Ala Leu Glu Met
                325                 330                 335

Val Tyr Thr Phe Gly Met Glu Asp Lys Phe Ser Ala Ala Leu Val Leu
            340                 345                 350

Thr Ser Phe Leu Lys Met Ser Lys Glu Ser Phe Glu Arg Ala Lys Arg
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Lys Ala Gln Ser Pro Leu Ala Phe Lys Glu Ala Ala Thr Lys Gln Leu
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Ala Val Leu Ser Ser Val Met Gln Cys Met Glu Thr His Lys Leu Asp
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Pro Ala Lys Glu Leu Pro Gly Trp Gln Ile Lys Glu Gln Ile Val Ser
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Leu Glu Lys Asp Thr Leu Gln Leu Asp Lys Glu Met Glu Glu Lys Ala
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Val Thr Ser Ser Ser Tyr Ser Pro Ile Tyr Arg Asp Arg Ser Phe Pro
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Ser Gln Arg Asp Asp Asp Gln Asp Glu Ile Ser Ala Leu Val Ser Ser
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Tyr Leu Gly Pro Ser Thr Ser Phe Pro His Arg Ser Arg Arg Ser Pro
            500                 505                 510

Glu Tyr Met Val Pro Leu Pro His Gly Gly Leu Gly Arg Ser Val Tyr
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Ala Tyr Glu His Leu Ala Pro Asn Ser Tyr Ser Pro Gly His Gly His
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Arg Leu His Arg Gln Tyr Ser Pro Ser Leu Val His Gly Gln Arg His
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Pro Leu Gln Tyr Ser Pro Pro Ile His Gly Gln Gln Gln Leu Pro Tyr
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Gly Ile Gln Arg Val Tyr Arg His Ser Pro Ser Glu Glu Arg Tyr Leu
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<223> Description of Artificial Sequence: Primer

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cDNA

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